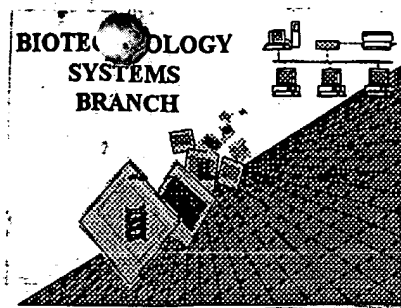


4a Counselor



#4
KAC
01-04-01

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/199,129
Source: 1635 RUSH
Date Processed by STIC: 1/4/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:45

Input Set : D:\slreg..txt
Output Set: N:\CRF3\01042001\I199129.raw

Does Not Comply
Corrected Diskette Needed

ppr 1-5

1 <110> APPLICANT: Byrum, Joseph R.
2 La Rosa, Thomas J.
4 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
5 Plants
7 <130> FILE REFERENCE: 38-21(15075)B
9 <140> CURRENT APPLICATION NUMBER: US/09/199,129
9 <141> CURRENT FILING DATE: 1998-11-24
9 <160> NUMBER OF SEQ ID NOS: 5521

OK

global errors

ERRORED SEQUENCES

11 <210> SEQ ID NO: 1
12 <211> LENGTH: 254
E--> 13 <212> TYPE: nucleic acid *invalid response per new*
14 <213> ORGANISM: Glycine max *Sequence Rules (1.823)*
16 <223> OTHER INFORMATION: Clone ID: 700547901H1 *→ L2207 (see below)*
18 <400> SEQUENCE: 1
20 ggaaatctga agctacctta agcgaatcn anagggccaa nngcaacac tagtcacgac 60
22 ttcaagctac ttgattttc cttdhacte tgcactcttc attcgctgag attttcttcg 120
24 aagagagatt cgaagatgga tgaagagacc gnggggtgtc atccgattgg aaaaggagtg 180
26 aaggaaaaagc ggttaccttc aagcacacag caggaaaatg gtctggggat ccgatgacn 240
28 aagggtcttca gaca *(see item 10 on error summary sheet)* 254
31 <210> SEQ ID NO: 2
32 <211> LENGTH: 285
E--> 33 <212> TYPE: nucleic acid
34 <213> ORGANISM: Glycine max *→ L2207*
36 <223> OTHER INFORMATION: Clone ID: 700547902H1
38 <400> SEQUENCE: 2
40 gggcgcttct gnggcgcgca cngetgttcc ttcgacgcgc ggcgngncgg gnggcgctgc 60
42 gccaaaccgg gactgcggcg ggtncctgt tctgcaacgg ctcggngga agccaccgn 120
44 ctagnctggc ggagtccacc ctgggcaacg ancatgantt ctacgaagt agcctggtgg 180
46 acgggtacnn cctaccatc tccatnanch ccttcaagg attccggaaa atgcactacn 240
48 ccgggttgcg tgaacgagnt caacnccatg tgcgccgttg ggcct 285
51 <210> SEQ ID NO: 3
52 <211> LENGTH: 277
E--> 53 <212> TYPE: nucleic acid
54 <213> ORGANISM: Glycine max *→ L2207*
56 <223> OTHER INFORMATION: Clone ID: 700547903H1
58 <400> SEQUENCE: 3
60 cgggtgaccca ttcattggaa gcttggaac ccaggttaca tccagccctt tgatcgcatg 60
62 gtaactgtcc aacctgccc catacagaac cgcagtyagc ccactactaa gagggatcga 120
64 ggtgggctcg gccacggat accttctggt gggccattc gtgaaggccg ggcctcttag 180
66 gaacaccgag atcgccgggc aagcgggctc tctggccgcc ggtgggcttg tggatgacct 240
68 cagcctttgc ctcacaatct ntgggatttc atccttc 277
71 <210> SEQ ID NO: 4
72 <211> LENGTH: 219

check for n/s →
in sequence

check for n/s

The only valid responses for a nucleotide sequence are either DNA or RNA. If the sequence is a combined DNA/RNA, use DNA and explain in L2207-L2237 section.

use L2207 (header only - no response) WHENEVER L2217, L2227, OR L2237 is shown. MANDATORY.

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set: N:\CRF3\01042001\I199129.raw

```

E--> 73 <212> TYPE: nucleic acid
      74 <213> ORGANISM: Glycine max
      76 <223> OTHER INFORMATION: Clone ID: 700547904H1
      78 <400> SEQUENCE: 4
      80 agccatcacc atgggagcaq tgaattctccc agatctcggc accgagattt tgattccggt 60
      82 ctgcgccacc atlggaatag ggltcgcctt ctccagtggt gtcctcgtct ccaagggtta 120
      84 gctctccgct gccagagacg ctccccctaa cgcgcgcggc aaaaatgqct acaacgatta 180
      86 cctcaccgna gaagaggaag gcctcaacga tcacnacgt 219
      89 <210> SEQ ID NO: 5
      90 <211> LENGTH: 271

E--> 91 <212> TYPE: nucleic acid
      92 <213> ORGANISM: Glycine max
      94 <223> OTHER INFORMATION: Clone ID: 700547905H1
      96 <400> SEQUENCE: 5
      98 ctgcgcacc ctlatcctcg ttgtctcttc ttlgcactc attcttggct ttttcgacga 60
      100 ctaccacct gcttcaaaag atgctacaaa ccggaagaag cactagtgtc atagttatca 120
      102 ttcttccagg ggataaaata ttaggctctg caaaaatttc ttgggtattg gaatcccagg 180
      104 agatgccaaq gacttcttca atcaagtcga gtcataaggt ttctagagag caataggaaa 240
      106 gtgcttacta aaaaacaatt tatttaaatg c 271
      109 <210> SEQ ID NO: 6
      110 <211> LENGTH: 245

E--> 111 <212> TYPE: nucleic acid
      112 <213> ORGANISM: Glycine max
      114 <223> OTHER INFORMATION: Clone ID: 700547906H1
      116 <400> SEQUENCE: 6
      118 gcggcttcca atggcaccac tgcaattcac ctgcaccga ctcaaccaca ttgtncagg 60
      120 ctctctacc ttccacgcga accttctctc gaaaccgaag cctcatttgc aaagccactt 120
      122 tcttcttgtt ctgacgactt ctccaaaagn aggcgcgnt tcagcgcn gn aancqccgna 180
      124 ctctcggnct cgtcccnca gttannngn cgtgncgaag angcgtttic ggagttggan 240
      126 anagt 245
      129 <210> SEQ ID NO: 7
      130 <211> LENGTH: 275

E--> 131 <212> TYPE: nucleic acid
      132 <213> ORGANISM: Glycine max
      134 <223> OTHER INFORMATION: Clone ID: 700547907H1
      136 <400> SEQUENCE: 7
      138 tcacaaaaca actttcaaac tctgagaaag aatggctgcc aacacattga tgagtactgc 60
      140 latctcagcc ttccactctc tctttcttct ctcaaaatcc agatttgcca ncgcagttcc 120
      142 tcttcttagc ttgggtgtca ccaatgcctc ttcttctcgc ttctctatga gtgctgactg 180
      144 gatgccagcg nagectagac ctccctlanct tgatgggtca gcacctgggt actttggatt 240
      146 cganccctctt ngctcttggtg aagtaccaga gaato 275
      149 <210> SEQ ID NO: 8
      150 <211> LENGTH: 273

E--> 151 <212> TYPE: nucleic acid
      152 <213> ORGANISM: Glycine max
      154 <223> OTHER INFORMATION: Clone ID: 700547909H1
      156 <400> SEQUENCE: 8
      158 anagcctagg gttcttttct tcttctgctc tcactccttg caaatctcgc gtaacgcatg 60
      160 cgtttcttca ctcaatcgga gccacgact gatctatttg gaactcttgc aggtctgttg 120

```

same
error

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set: N:\CRF3\01042001\I199129.raw

```

162 ttttactgta ctcaatttga taaatraagg cccccccccn nnnnnnnnnn nnnntcccc 180
164 tgggtgttlat ggatgtgtcc atcgatgggg atcctgttga aaggalggtt tttagacttt 240
166 ctatgatgtt qctcccaaga ctgnagaaaa ctt 273
169 <210> SEQ ID NO: 9
170 <211> LENGTH: 235
E--> 171 <212> TYPE: nucleic acid
172 <213> ORGANISM: Glycine max
174 <223> OTHER INFORMATION: Clone ID: 700547910H1
176 <400> SEQUENCE: 9
178 agcccatcca aggataacga atttatgggt ggatcaattt caaacaggag gcgttctagg 60
180 caacgttcgg cctcaaatit tcttctcgg tatcaacaac ctcatctccc acaagccaa 120
182 gataatgggt ctgtggggca ttatgggtat tcattctcga qctatagttg tggctgtgt 180
184 ccagaacaag taaagugctn ggataggaag tatttaaggg taggtgatga ttata 235
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 265
E--> 189 <212> TYPE: nucleic acid
190 <213> ORGANISM: Glycine max
192 <223> OTHER INFORMATION: Clone ID: 700547911H1
194 <400> SEQUENCE: 10
196 caaaaaaccc aaatgcaggt tcttcacta aaacttgggc qattttcana ttngtactt 60
198 tggcaccctt cactagtag ttctaccat gtccgatact aaaagaagt ccaacaacag 120
200 cggttccaag cggagcctgc catcgtggac gaattcaagg qaaaatgaga gcgataacag 180
202 tgcaaaagaa ccaactttgg atggccaagg tgagaaatcc agtgacgctg agacacccca 240
204 caagagcaaa gtccaaaatg aaat 265
207 <210> SEQ ID NO: 11
208 <211> LENGTH: 273
E--> 209 <212> TYPE: nucleic acid
210 <213> ORGANISM: Glycine max
212 <223> OTHER INFORMATION: Clone ID: 700547913H1
214 <400> SEQUENCE: 11
216 gtcacaaccc aagcctccgc cgcgatttct cggccatgtg cctcgaaatc gaggtttctg 60
218 accgctcttt ccggtaaaact caaccyggaa gtgactatga gqccaatggg gtgccctcct 120
220 tctgcctctt tcaagggttg agccaagaag ggagagtggg tacctggctt ggccctccca 180
222 acttacctca atggcaactt tctgtgtgac aatggatttg accctctggg actagctgag 240
224 gaccagaga acttgaggtg gtacgttcaa gcc 273
227 <210> SEQ ID NO: 12
228 <211> LENGTH: 273
E--> 229 <212> TYPE: nucleic acid
230 <213> ORGANISM: Glycine max
232 <223> OTHER INFORMATION: Clone ID: 700547914H1
234 <400> SEQUENCE: 12
236 gcgtacaac aataaaaatc tccatttgt nttttcttct cttaacgcac accaacaana 60
238 cctctctctc atcggaatct ccaaaaagaa tggacaacaa aacgcagcag tccgagagta 120
240 agcaaaaacga caacgacgag gaagttgcgc caaaacgaca agaccctaac ccgtcgtctg 180
242 gcgggtgggg cttttcaccg ctctcgttct tctccgatct tcagaaggcc gccgctgttg 240
244 cagccgaaga gatctctcgc aatgtgctg tag 273
247 <210> SEQ ID NO: 13
248 <211> LENGTH: 277
E--> 249 <212> TYPE: nucleic acid

```

same

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\sireg..txt
 Output Set : N:\CRF3\01042001\I199129.raw

```

250 <213> ORGANISM: Glycine max
252 <223> OTHER INFORMATION: Clone ID: 700547916H1
254 <400> SEQUENCE: 13
256 agaacaagta gttgaqaact aagaagqaga agcaaatggc ttcctcaatg atctcttccc 60
258 cagctgttac cactgtcaac cgtgcegggt cgggcatggt tgetccatc actggcctca 120
260 agtccatggc tggcttcccc accagaaaga ccaacaatga cattanctcc attgctagca 180
262 acggtggaag agtgcnatgc atgcaqgtgt gggccaccag ttggcaagaa gaagtttgag 240
264 actcttccct acctgccaga ccttgatgat gcacatt 277
267 <210> SEQ ID NO: 14
268 <211> LENGTH: 275
E--> 269 <212> TYPE: nucleic acid
270 <213> ORGANISM: Glycine max
272 <223> OTHER INFORMATION: Clone ID: 700547917H1
274 <400> SEQUENCE: 14
276 qagcatttcn aatggagcaa gaaatttqgt attcatttct tttcgttggg nattgcattn 60
278 qcaatgattc agattcattg attgcttgc nacgggnct tgggttctac tctgaacacn 120
280 aattttcaag atagctatgg gtgttgnata tttttntgtg aagaattaat gagcataatg 180
282 ttttctggaa agantggggg cagacgaatc tcattgtcaa ganttacaga gtttgagtgt 240
284 gtcaaaagca cttgtganga gtgttagcca gaagt 275
287 <210> SEQ ID NO: 15
288 <211> LENGTH: 273
E--> 289 <212> TYPE: nucleic acid
290 <213> ORGANISM: Glycine max
292 <223> OTHER INFORMATION: Clone ID: 700547920H1
294 <400> SEQUENCE: 15
296 attctctcca tatattatct caaacccctc tcacagaatg ggaagtgtg gaggaactga 60
298 ctatgttgca tacacttatg agaattctga gagagagcct tactggccat cagagaagct 120
300 taagatttcc atcactggng ctgggggttt tatcgcgtca cacatagctc ggcgcctcaa 180
302 gacagagggg cattacatta ttgcttctga ttggaagaaa aatgagcaca tgactyagga 240
304 catgttctgt gatgaattcc atcttgttga tct 273
307 <210> SEQ ID NO: 16
308 <211> LENGTH: 273
E--> 309 <212> TYPE: nucleic acid
310 <213> ORGANISM: Glycine max
312 <223> OTHER INFORMATION: Clone ID: 700547921H1
314 <400> SEQUENCE: 16
316 atcacaccat gccacgcta gtyacaatca agagaactcc ttcaagaagg gttctgctcg 60
318 ctctgccagt aaaagtcagg agaacaagtc atctggatta tcaaaagcat cgactaatgc 120
320 aaataattat gggtcagttt ctctctcaag ntcaagtgtc cctgcaaaac gtactgagga 180
322 tgacatggat gattttgatc caagagggaac ttctaccaa acttcagctg gaaactctaa 240
324 cccagattga tctcttttga caagattaat cgg 273
327 <210> SEQ ID NO: 17
328 <211> LENGTH: 245
E--> 329 <212> TYPE: nucleic acid
330 <213> ORGANISM: Glycine max
332 <223> OTHER INFORMATION: Clone ID: 700547924H1
334 <400> SEQUENCE: 17
336 gacacgtca cgcctgcaaa ccaacttgag tgcgttccag gaaacacatc cctctccat 60
338 ttcatcact tctctgctct ctctttctcc cactgaattcc caaacttatt ttcacgttc 120

```

Same

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/199,129
 DATE: 01/04/2001
 TIME: 11:43:45

Input Set : D:\slreg..txt
 Output Set: N:\CRF3\01042001\I199129.raw

```

340 caattttccgt cagctaaacc atgcataacc gectcccaaa ttaggggttc cgacaggcgc 180
342 gtggagggaat cgtatggcgc cgtgaatccg cagcctctgc agttcgagga ccttgcctata 240
344 cccgt 245
347 <210> SEQ ID NO: 18
348 <211> LENGTH: 271
E--> 349 <212> TYPE: nucleic acid
350 <213> ORGANISM: Glycine max
352 <223> OTHER INFORMATION: Clone ID: 700547925H1
354 <400> SEQUENCE: 18
356 ttcttccca ttccaccct cgtcttccct tgtttccggt gagggccaac caaacatat 60
358 gttaaacag gaacacacct caaacataa catcctacgc caaacaaaac ttgacctcc 120
360 ttcagatcta taccctnnnn nnnnnnnnn ntctattttc aaattctaca tcatgggcac 180
362 qgaggttcta cggccacaag attgtttcac ccaacgcac ggtgttccac cgcctggctt 240
364 ttcccggcga agaacctatg gtaccacca c 271
367 <210> SEQ ID NO: 19
368 <211> LENGTH: 270
E--> 369 <212> TYPE: nucleic acid
370 <213> ORGANISM: Glycine max
372 <223> OTHER INFORMATION: Clone ID: 700547926H1
374 <400> SEQUENCE: 19
376 ccttatitc ttccctagt caaggagatt catctctgca gcagggtgtt ttagaaaaga 60
378 aaaaatgclg aaagtltgct gcattggggc tggatatgtg ggggttccaa caatggcagt 120
380 cattgcactg aaatgccct caattgaagt ggtgtgtgtt gacatctcta aatccaggat 180
382 tgcagcatgg aacagtgaac agctccctat ctatgagcca ggccttgatg atgtggtgaa 240
384 qcaatgtcgt qgcaagaacc tttcttcag 270
387 <210> SEQ ID NO: 20
388 <211> LENGTH: 269
E--> 389 <212> TYPE: nucleic acid
390 <213> ORGANISM: Glycine max
392 <223> OTHER INFORMATION: Clone ID: 700547927H1
394 <400> SEQUENCE: 20
396 cnaqatttga aggttcgtta tgcctccat ctcccaactg tcttgccgcg tcttacatgc 60
398 aaatttcgtg gaggactgaa aactggcatt gttgggagaa cagggaagtgg taaatccact 120
400 ctcatacaaa cacttttccg aattgttgag cctactgccg gccaaqttat gattgacagc 180
402 atcaacatct ctccaatgga ctccatgatt tgaggtctag actaagcacc atcctccaga 240
404 ttccacaatg ttgaaggga cgtgagaa 269

```

same

FYI:

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

FYI:

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:57

Input Set : D:\slreg..txt
Output Set: N:\CRF3\01042001\I199129.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:13 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:33 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:53 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:73 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:91 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:111 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:131 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:151 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:171 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:189 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:209 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:229 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:249 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:269 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:289 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:309 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:329 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:349 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:369 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:389 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:409 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:429 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:449 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:469 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:489 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:509 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:529 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:549 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:569 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:589 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:609 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:629 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:649 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:669 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:689 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:709 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:729 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:749 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:769 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:789 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:809 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:829 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:849 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:863 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:883 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:903 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/199,129

DATE: 01/04/2001
TIME: 11:43:57

Input Set : D:\slreg..txt
Output Set: N:\CRF3\01042001\I199129.raw

L:923 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:943 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:963 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:983 M:310 E: (3) Wrong or Missing Sequence Type, TYPE: